	Application No.	Application No. Applicant(s)		
Notice of Allowability	08/170,344	KAST ET AL.		
	Examiner	Art Unit		
	N. M. Minnifield	1645		
The MAILING DATE of this communication apper All claims being allowable, PROSECUTION ON THE MERITS IS herewith (or previously mailed), a Notice of Allowance (PTOL-85) NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RI of the Office or upon petition by the applicant. See 37 CFR 1.313	(OR REMAINS) CLOSED in t or other appropriate commun GHTS. This application is sui	his application. If not includication will be mailed in due	led course. THIS	
1. This communication is responsive to 11/20/06; 12/6/07.				
2. The allowed claim(s) is/are 5, 6, 8, 10, 12, 14, 16 and 26-3	0; now renumbered 1-12 resp	ectively.		
 3. Acknowledgment is made of a claim for foreign priority una) All b) Some* c) None of the: 1. Certified copies of the priority documents have 2. Certified copies of the priority documents have 3. Copies of the certified copies of the priority documents have International Bureau (PCT Rule 17.2(a)). * Certified copies not received: 	been received. been received in Application	No	ation from the	
Applicant has THREE MONTHS FROM THE "MAILING DATE" noted below. Failure to timely comply will result in ABANDONM THIS THREE-MONTH PERIOD IS NOT EXTENDABLE.		reply complying with the re	equirements	
4. A SUBSTITUTE OATH OR DECLARATION must be subminformal PATENT APPLICATION (PTO-152) which give			NOTICE OF	
5. X CORRECTED DRAWINGS (as "replacement sheets") mus	t be submitted.	·		
(a) ☐ including changes required by the Notice of Draftspers		PTO-948) attached		
1) ☐ hereto or 2) ☑ to Paper No./Mail Date <u>#13/08</u>	3 <u>-23-95</u> .			
(b) including changes required by the attached Examiner's Paper No./Mail Date	s Amendment / Comment or in	n the Office action of		
Identifying indicia such as the application number (see 37 CFR 1. each sheet. Replacement sheet(s) should be labeled as such in the			e back) of	
6. DEPOSIT OF and/or INFORMATION about the deposit attached Examiner's comment regarding REQUIREMENT I			Note the	
Attachment(s)				
1. Notice of References Cited (PTO-892)		rmal Patent Application		
2. Notice of Draftperson's Patent Drawing Review (PTO-948)		 Interview Summary (PTO-413), Paper No./Mail Date <u>attached</u>. 		
 Information Disclosure Statements (PTO/SB/08), Paper No./Mail Date Examiner's Comment Regarding Requirement for Deposit of Biological Material 	7. X Examiner's Amendment/Comment			
		8. Examiner's Statement of Reasons for Allowance9. Other		
		N. M. Minnifield Primary Examiner Art Unit: 1645		

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EXAMINER'S AMENDMENT

- 1. Applicants' amendments filed November 20, 2006 and September 19, 2007 are acknowledged and have been entered. Claims 1-4, 7, 9, 11, 13, 15 and 17-25 have been canceled. Claims 5, 6, 8, 10, 12, 14, 16 and 26-30 have been amended. Claims 5, 6, 8, 10, 12, 14, 16 and 26-30 are now pending in the present application. All rejections/objections have been withdrawn in view of Applicants' amendment to the claims and/or comments.
- 2. An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this examiner's amendment was given in a telephone interview with Mita Mukherjee, 54325, on December 6, 2007.

- 3. The application has been amended as follows:
 - 1-4. (cancelled)
- 5. (currently amended) [A] <u>An</u> isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7- 15 of HPV16 protein E6) SEQ ID NO:1 KLPQLCTEL (residues 18- 26 of HPV16 protein E6) SEQ ID NO:2 QLCTELQTT (residues 21- 29 of HPV16 protein E6) SEQ ID NO:3 LCTELQTTI (residues 22- 30 of HPV16 protein E6) SEQ ID NO:4

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ELQTTIHDI (residues 25-33 of HPV16 protein E6) SEQ ID NO:5 LQTTIHDII (residues 26-34 of HPV16 protein E6) SEQ ID NO:6 TIHDIILEC (residues 29- 37 of HPV16 protein E6) SEQ ID NO:7 IHDIILECV (residues 30-38 of HPV16 protein E6) SEQ ID NO:8 CVYCKQQLL (residues 37- 45 of HPV16 protein E6) SEQ ID NO:9 FAFRDLCIV (residues 52-60 of HPV16 protein E6) SEO ID NO:10 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 PLCDLLIRC (residues 102-110 of HPV16 protein E6) SEQ ID NO:12 TLHEYMLDL (residues 7-15 of HPV16 protein E7) SEQ ID NO:13 MLDLQPETT (residues 12-20 of HPV16 protein E7) SEQ ID NO:15 RLCVQSTHV (residues 66-74 of HPV16 protein E7) SEQ ID NO:16 TLEDLLMGT (residues 78-86 of HPV16 protein E7) SEQ ID NO:17 LLMGTLGIV (residues 82-90 of HPV16 protein E7) SEQ ID NO:18 GTLGIVCPI (residues 85-93 of HPV16 protein E7) SEQ ID NO:19 and TLGIVCPIC (residues 86-94 of HPV16 protein E7) SEQ ID NO:20; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A2.1.

6. (previously presented) An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of: KLPDLCTEL (residues 13- 21 of HPV18 protein E6) SEQ ID NO:21 SLQDIEITC (residues 24- 32 of HPV18 protein E6) SEQ ID NO:22

LQDIEITCV (residues 25-33 of HPV18 protein E6) SEQ ID NO:23

EITCVYCKT (residues 29- 37 of HPV18 protein E6) SEQ ID NO:24

KTVLELTEV (residues 36-44 of HPV18 protein E6) SEQ ID NO:25

ELTEVFEFA (residues 40-48 of HPV18 protein E6) SEQ ID NO:26

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FAFKDLFVV (residues 47- 55 of HPV18 protein E6) SEQ ID NO:27
DTLEKLTNT (residues 88- 96 of HPV18 protein E6) SEQ ID NO:28
LTNTGLYNL (residues 93-101 of HPV18 protein E6) SEQ ID NO:29
TLQDIVLHL (residues 7- 15 of HPV18 protein E7) SEQ ID NO:30
FQQLFLNTL (residues 86- 94 of HPV18 protein E7) SEQ ID NO:31
QLFLNTLSF (residues 88- 96 of HPV18 protein E7) SEQ ID NO:32
LFLNTLSFV (residues 89- 97 of HPV18 protein E7) SEQ ID NO:33 and
LSFVCPWCA (residues 94-102 of HPV18 protein E7) SEQ ID NO:34;
wherein said amino acid sequence is derived from protein E6 or E7 of HPV18; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A2.1.

7. (cancelled)

8. (currently amended) [A] <u>An isolated</u> peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

YRDGNPYAV (residues 61- 69 of HPV16 protein E6) SEQ ID NO:35
WTGRCMSCC (residues 139-147 of HPV16 protein E6) SEQ ID NO:36
MSCCRSSRT (residues 144-152 of HPV16 protein E6) SEQ ID NO:37
TTDLYCYEQ (residues 19- 27 of HPV16 protein E7) SEQ ID NO:38
EIDGPAGQA (residues 37- 45 of HPV16 protein E7) SEQ ID NO:39 and
HVDIRTLED (residues 73- 81 of HPV16 protein E7) SEQ ID NO:40;
wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A1.

9. (cancelled)

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10. (currently amended) [A] <u>An isolated</u> peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7-15 of HPV16 protein E6) SEQ ID NO:1 IILECVYCK (residues 33-41 of HPV16 protein E6) SEQ ID NO:41 CVYCKQQLL (residues 37-45 of HPV16 protein E6) SEQ ID NO:9 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 QQLLRREVY (residues 42-50 of HPV16 protein E6) SEQ ID NO:43 IVYRDGNPY (residues 59-67 of HPV16 protein E6) SEQ ID NO:44 YAVCDKCLK (residues 67-75 of HPV16 protein E6) SEQ ID NO:45 AVCDKCLKF (residues 68-76 of HPV16 protein E6) SEQ ID NO:46 VCDKCLKFY (residues 69-77 of HPV16 protein E6) SEQ ID NO:47 KFYSKISEY (residues 75-83 of HPV16 protein E6) SEQ ID NO:48 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 ISEYRHYCY (residues 80-88 of HPV16 protein E6) SEQ ID NO:49 RHYCYSLYG (residues 84-92 of HPV16 protein E6) SEQ ID NO:50 SLYGTTLEQ (residues 89-97 of HPV16 protein E6) SEQ ID NO:51 TTLEQQYNK (residues 93-101 of HPV16 protein E6) SEQ ID NO:52 QQYNKPLCD (residues 97-105 of HPV16 protein E6) SEQ ID NO:53 LIRCINCQK (residues 107-115 of HPV16 protein E6) SEQ ID NO:54 HLDKKQRFH (residues 125-133 of HPV16 protein E6) SEQ ID NO:55 CMSCCRSSR (residues 143-151 of HPV16 protein E6) SEQ ID NO:56 SCCRSSRTR (residues 145-153 of HPV16 protein E6) SEQ ID NO:57 CCRSSRTRR (residues 146-154 of HPV16 protein E6) SEQ ID NO:58 YNIVTFCCK (residues 52-60 of HPV16 protein E7) SEQ ID NO:60 CCKCDSTLR (residues 58- 66 of HPV16 protein E7) SEQ ID NO:61 and KCDSTLRLC (residues 60- 68 of HPV16 protein E7) SEQ ID NO:62;

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wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A3.2.

11. (cancelled)

12. (currently amended) [A] <u>An isolated</u> peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7-15 of HPV16 protein E6) SEQ ID NO:1 IILECVYCK (residues 33-41 of HPV16 protein E6) SEQ ID NO:41 CVYCKQQLL (residues 37-45 of HPV16 protein E6) SEQ ID NO:9 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 QQLLRREVY (residues 42-50 of HPV16 protein E6) SEQ ID NO:43 IVYRDGNPY (residues 59-67 of HPV16 protein E6) SEQ ID NO:44 YAVCDKCLK (residues 67-75 of HPV16 protein E6) SEQ ID NO:45 AVCDKCLKF (residues 68-76 of HPV16 protein E6) SEQ ID NO:46 VCDKCLKFY (residues 69-77 of HPV16 protein E6) SEO ID NO:47 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 ISEYRHYCY (residues 80-88 of HPV16 protein E6) SEQ ID NO:49 LIRCINCOK (residues 107-115 of HPV16 protein E6) SEO ID NO:54 TGRCMSCCR (residues 140-148 of HPV16 protein E6) SEQ ID NO:63 CMSCCRSSR (residues 143-151 of HPV16 protein E6) SEQ ID NO:56 SCCRSSRTR (residues 145-153 of HPV16 protein E6) SEQ ID NO:57 YNIVTFCCK (residues 52-60 of HPV16 protein E7) SEQ ID NO:60 CCKCDSTLR.(residues 58-66 of HPV16 protein E7) SEQ ID NO:61 and VCPICSQKP (residues 90- 98 of HPV16 protein E7) SEQ ID NO:64; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and

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wherein said peptide has the ability to bind to human MHC Class I allele HLA-A11.2.

(cancelled) 13.

(currently amended) [A] An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

MHQKRTAMF (residues 1-9 of HPV16 protein E6) SEQ ID NO:65 LQTTIHDII (residues 26- 34 of HPV16 protein E6) SEQ ID NO:6 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 LLRREVYDF (residues 44-52 of HPV16 protein E6) SEQ ID NO:66 VYDFAFRDL (residues 49-57 of HPV16 protein E6) SEQ ID NO:67 PYAVCDKCL (residues 66-74 of HPV16protein E6) SEQ ID NO:68 KCLKFYSKI (residues 72-80 of HPV16 protein E6) SEQ ID NO:69 EYRHYCYSL (residues 82-90 of HPV16 protein E6) SEQ ID NO:70 HYCYSLYGT (residues 85-93 of HPV16 protein E6) SEQ ID NO:71 CYSLYGTTL (residues 87-95 of HPV16 protein E6) SEQ ID NO:72 RFHNIRGRW (residues 131-139 of HPV16 protein E6) SEQ ID NO:73 and RAHYNIVTF (residues 49-57 of HPV16 protein E7) SEQ ID NO:74; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-

(cancelled) 15.

A24.

(previously presented) A pharmaceutical composition comprising the 16. peptide of claim 5 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.

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17-25. (cancelled)

- 26. (previously presented) A pharmaceutical composition comprising the peptide of claim 6 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 27. (previously presented) A pharmaceutical composition comprising the peptide of claim 8 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 28. (previously presented) A pharmaceutical composition comprising the peptide of claim 10 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 29. (previously presented) A pharmaceutical composition comprising the peptide of claim 12 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 30. (previously presented) A pharmaceutical composition comprising the peptide of claim 14 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 4. Claims 5, 6, 8, 10, 12, 14, 16 and 26-30 have been allowed and renumbered 1-12 respectively.
- 5. Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee.

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6. Any inquiry concerning this communication or earlier communications from the examiner should be directed to N. M. Minnifield whose telephone number is 571-272-0860. The examiner can normally be reached on M-F (8:00-5:30) Second Friday Off.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Shanon Foley can be reached on 571-272-8975. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

Primary Examiner

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NMM

December 7, 2007

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CLEAN COPY OF ALLOWED CLAIMS

5. An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7-15 of HPV16 protein E6) SEQ ID NO:1 KLPOLCTEL (residues 18-26 of HPV16 protein E6) SEQ ID NO:2 QLCTELQTT (residues 21-29 of HPV16 protein E6) SEQ ID NO:3 LCTELQTTI (residues 22-30 of HPV16 protein E6) SEQ ID NO:4 ELOTTIHDI (residues 25-33 of HPV16 protein E6) SEQ ID NO:5 LQTTIHDII (residues 26-34 of HPV16 protein E6) SEQ ID NO:6 TIHDIILEC (residues 29- 37 of HPV16 protein E6) SEQ ID NO:7 IHDIILECV (residues 30-38 of HPV16 protein E6) SEQ ID NO:8 CVYCKOOLL (residues 37-45 of HPV16 protein E6) SEQ ID NO:9 FAFRDLCIV (residues 52-60 of HPV16 protein E6) SEQ ID NO:10 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 PLCDLLIRC (residues 102-110 of HPV16 protein E6) SEQ ID NO:12 TLHEYMLDL (residues 7-15 of HPV16 protein E7) SEQ ID NO:13 MLDLQPETT (residues 12-20 of HPV16 protein E7) SEQ ID NO:15 RLCVQSTHV (residues 66-74 of HPV16 protein E7) SEQ ID NO:16 TLEDLLMGT (residues 78-86 of HPV16 protein E7) SEQ ID NO:17 LLMGTLGIV (residues 82-90 of HPV16 protein E7) SEQ ID NO:18 GTLGIVCPI (residues 85-93 of HPV16 protein E7) SEQ ID NO:19 and TLGIVCPIC (residues 86- 94 of HPV16 protein E7) SEQ ID NO:20;

wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A2.1.

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A2.1.

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An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

KLPDLCTEL (residues 13-21 of HPV18 protein E6) SEQ ID NO:21 SLQDIEITC (residues 24-32 of HPV18 protein E6) SEQ ID NO:22 LQDIEITCV (residues 25-33 of HPV18 protein E6) SEQ ID NO:23 EITCVYCKT (residues 29-37 of HPV18 protein E6) SEQ ID NO:24 KTVLELTEV (residues 36-44 of HPV18 protein E6) SEQ ID NO:25 ELTEVFEFA (residues 40-48 of HPV18 protein E6) SEQ ID NO:26 FAFKDLFVV (residues 47-55 of HPV18 protein E6) SEQ ID NO:27 DTLEKLTNT (residues 88-96 of HPV18 protein E6) SEQ ID NO:28 LTNTGLYNL (residues 93-101 of HPV18 protein E6) SEQ ID NO:29 TLQDIVLHL (residues 7-15 of HPV18 protein E7) SEQ ID NO:30 FQQLFLNTL (residues 86-94 of HPV18 protein E7) SEQ ID NO:31 OLFLNTLSF (residues 88-96 of HPV18 protein E7) SEQ ID NO:32 LFLNTLSFV (residues 89-97 of HPV18 protein E7) SEQ ID NO:33 and LSFVCPWCA (residues 94-102 of HPV18 protein E7) SEQ ID NO:34; wherein said amino acid sequence is derived from protein E6 or E7 of HPV18; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-

An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

YRDGNPYAV (residues 61-69 of HPV16 protein E6) SEQ ID NO:35 WTGRCMSCC (residues 139-147 of HPV16 protein E6) SEQ ID NO:36 MSCCRSSRT (residues 144-152 of HPV16 protein E6) SEQ ID NO:37 TTDLYCYEQ (residues 19-27 of HPV16 protein E7) SEQ ID NO:38 EIDGPAGQA (residues 37-45 of HPV16 protein E7) SEQ ID NO:39 and HVDIRTLED (residues 73-81 of HPV16 protein E7) SEQ ID NO:40;

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wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A1.

10. An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7-15 of HPV16 protein E6) SEQ ID NO:1 IILECVYCK (residues 33-41 of HPV16 protein E6) SEQ ID NO:41 CVYCKQQLL (residues 37-45 of HPV16 protein E6) SEQ ID NO:9 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 OOLLRREVY (residues 42-50 of HPV16 protein E6) SEQ ID NO:43 IVYRDGNPY (residues 59-67 of HPV16 protein E6) SEQ ID NO:44 YAVCDKCLK (residues 67-75 of HPV16 protein E6) SEQ ID NO:45 AVCDKCLKF (residues 68-76 of HPV16 protein E6) SEQ ID NO:46 VCDKCLKFY (residues 69-77 of HPV16 protein E6) SEQ ID NO:47 KFYSKISEY (residues 75-83 of HPV16 protein E6) SEQ ID NO:48 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 ISEYRHYCY (residues 80-88 of HPV16 protein E6) SEQ ID NO:49 RHYCYSLYG (residues 84-92 of HPV16 protein E6) SEQ ID NO:50 SLYGTTLEQ (residues 89-97 of HPV16 protein E6) SEQ ID NO:51 TTLEQQYNK (residues 93-101 of HPV16 protein E6) SEQ ID NO:52 QQYNKPLCD (residues 97-105 of HPV16 protein E6) SEQ ID NO:53 LIRCINCOK (residues 107-115 of HPV16 protein E6) SEQ ID NO:54 HLDKKQRFH (residues 125-133 of HPV16 protein E6) SEQ ID NO:55 CMSCCRSSR (residues 143-151 of HPV16 protein E6) SEQ ID NO:56 SCCRSSRTR (residues 145-153 of HPV16 protein E6) SEQ ID NO:57 CCRSSRTRR (residues 146-154 of HPV16 protein E6) SEQ ID NO:58 YNIVTFCCK (residues 52-60 of HPV16 protein E7) SEQ ID NO:60

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CCKCDSTLR (residues 58- 66 of HPV16 protein E7) SEQ ID NO:61 and KCDSTLRLC (residues 60- 68 of HPV16 protein E7) SEQ ID NO:62; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-A3.2.

12. An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

AMFQDPQER (residues 7-15 of HPV16 protein E6) SEQ ID NO:1 IILECVYCK (residues 33-41 of HPV16 protein E6) SEQ ID NO:41 CVYCKOOLL (residues 37- 45 of HPV16 protein E6) SEQ ID NO:9 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 OOLLRREVY (residues 42-50 of HPV16 protein E6) SEQ ID NO:43 IVYRDGNPY (residues 59-67 of HPV16 protein E6) SEQ ID NO:44 YAVCDKCLK (residues 67-75 of HPV16 protein E6) SEQ ID NO:45 AVCDKCLKF (residues 68-76 of HPV16 protein E6) SEQ ID NO:46 VCDKCLKFY (residues 69-77 of HPV16 protein E6) SEQ ID NO:47 KISEYRHYC (residues 79-87 of HPV16 protein E6) SEQ ID NO:11 ISEYRHYCY (residues 80-88 of HPV16 protein E6) SEQ ID NO:49 LIRCINCQK (residues 107-115 of HPV16 protein E6) SEQ ID NO:54 TGRCMSCCR (residues 140-148 of HPV16 protein E6) SEQ ID NO:63 CMSCCRSSR (residues 143-151 of HPV16 protein E6) SEQ ID NO:56 SCCRSSRTR (residues 145-153 of HPV16 protein E6) SEQ ID NO:57 YNIVTFCCK (residues 52-60 of HPV16 protein E7) SEQ ID NO:60 CCKCDSTLR.(residues 58-66 of HPV16 protein E7) SEQ ID NO:61 and VCPICSOKP (residues 90- 98 of HPV16 protein E7) SEQ ID NO:64; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and

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wherein said peptide has the ability to bind to human MHC Class I allele HLA-A11.2.

An isolated peptide comprising an amino acid sequence from 9 to 12 amino acids in length selected from the group consisting of:

MHQKRTAMF (residues 1-9 of HPV16 protein E6) SEQ ID NO:65 LQTTIHDII (residues 26- 34 of HPV16 protein E6) SEQ ID NO:6 VYCKQQLLR (residues 38-46 of HPV16 protein E6) SEQ ID NO:42 LLRREVYDF (residues 44-52 of HPV16 protein E6) SEQ ID NO:66 VYDFAFRDL (residues 49-57 of HPV16 protein E6) SEQ ID NO:67 PYAVCDKCL (residues 66-74 of HPV16protein E6) SEQ ID NO:68 KCLKFYSKI (residues 72-80 of HPV16 protein E6) SEQ ID NO:69 EYRHYCYSL (residues 82-90 of HPV16 protein E6) SEQ ID NO:70 HYCYSLYGT (residues 85-93 of HPV16 protein E6) SEQ ID NO:71 CYSLYGTTL (residues 87-95 of HPV16 protein E6) SEQ ID NO:72 RFHNIRGRW (residues 131-139 of HPV16 protein E6) SEQ ID NO:73 and RAHYNIVTF (residues 49- 57 of HPV16 protein E7) SEQ ID NO:74; wherein said amino acid sequence is derived from protein E6 or E7 of HPV16; and wherein said peptide has the ability to bind to human MHC Class I allele HLA-

- A pharmaceutical composition comprising the peptide of claim 5 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- A pharmaceutical composition comprising the peptide of claim 6 and 26. a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- A pharmaceutical composition comprising the peptide of claim 8 and 27. a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.

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- 28. A pharmaceutical composition comprising the peptide of claim 10 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 29. A pharmaceutical composition comprising the peptide of claim 12 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.
- 30. A pharmaceutical composition comprising the peptide of claim 14 and a pharmaceutically acceptable carrier, diluent, excipient or adjuvant.